

FIGURE 1

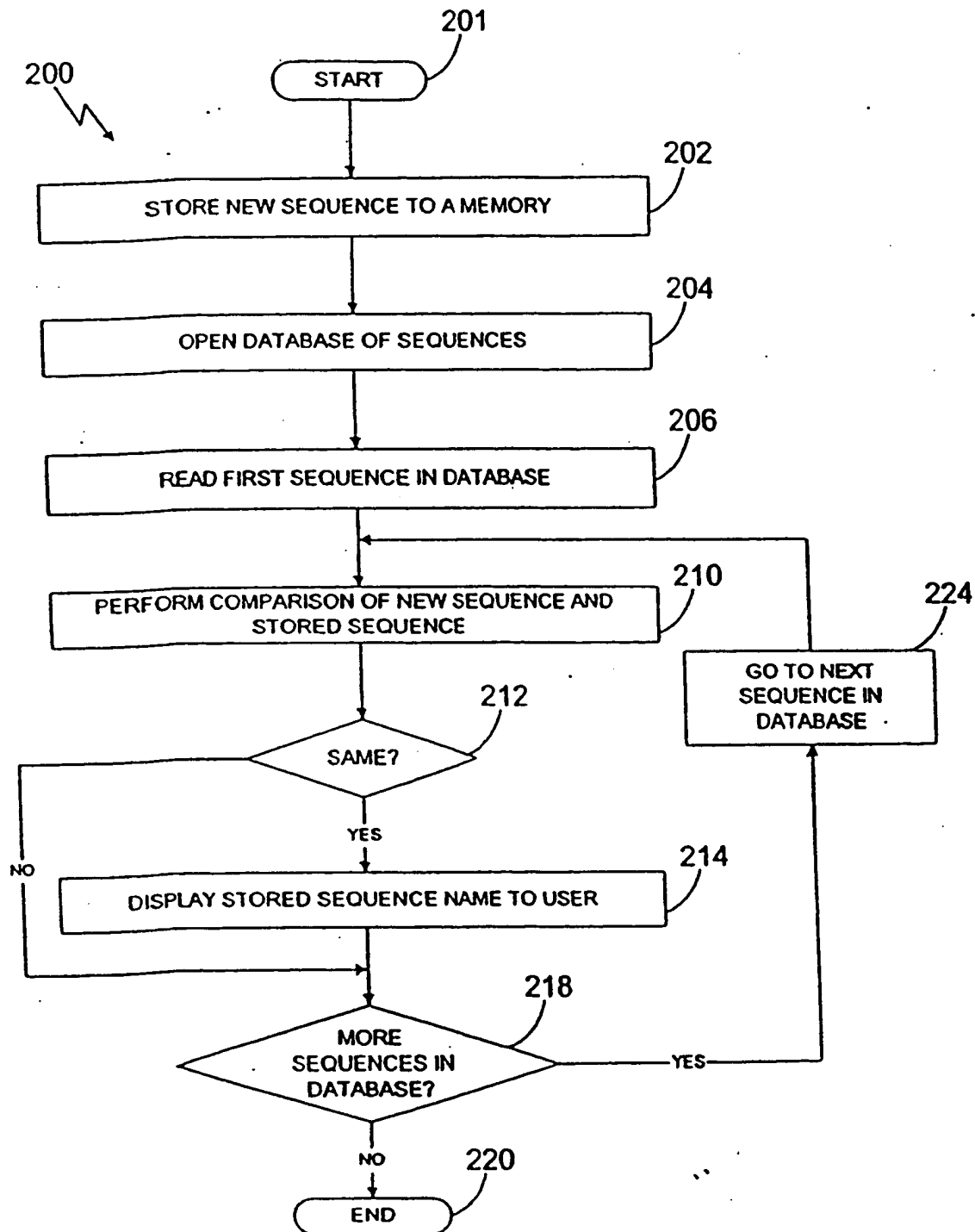


FIGURE 2

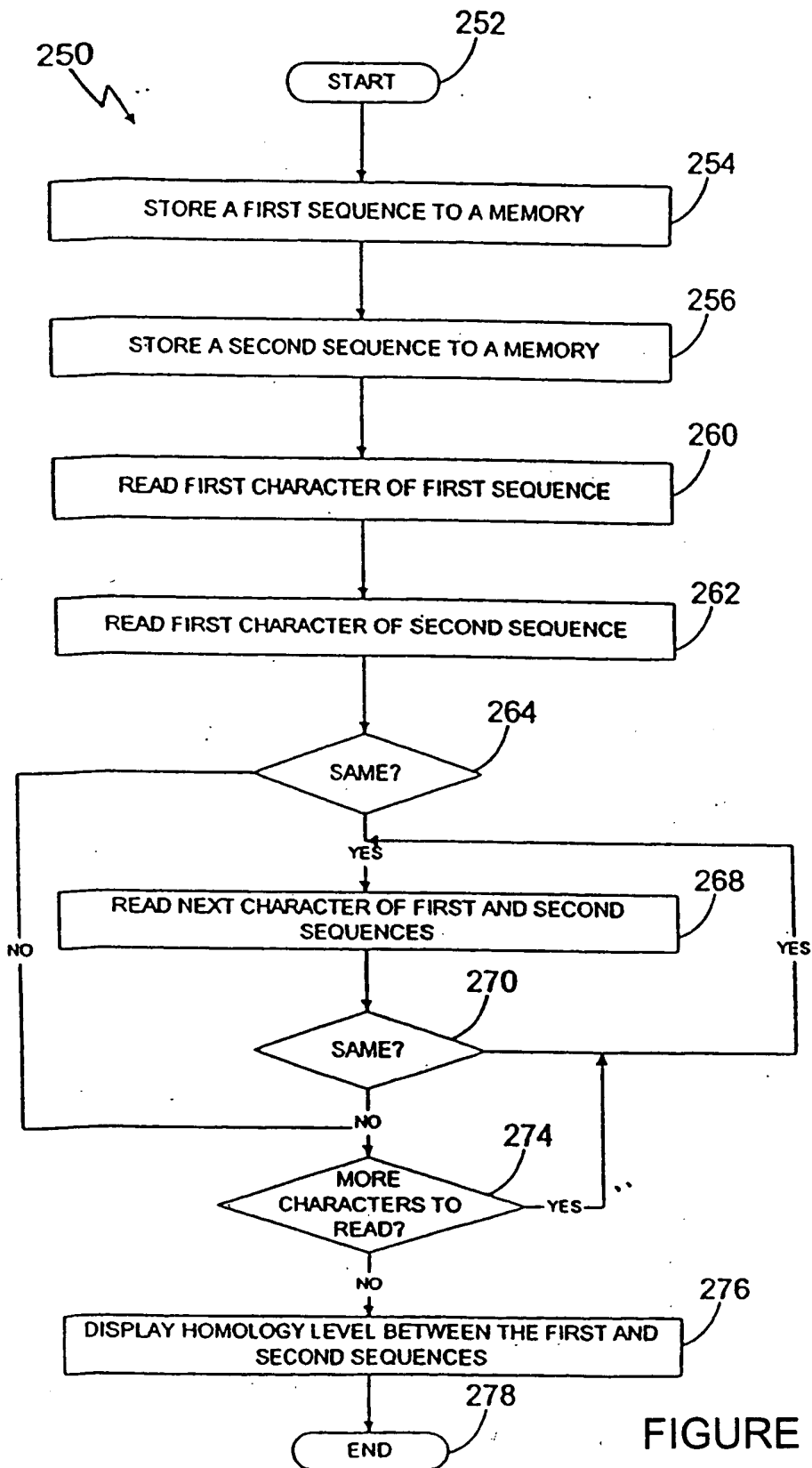


FIGURE 3

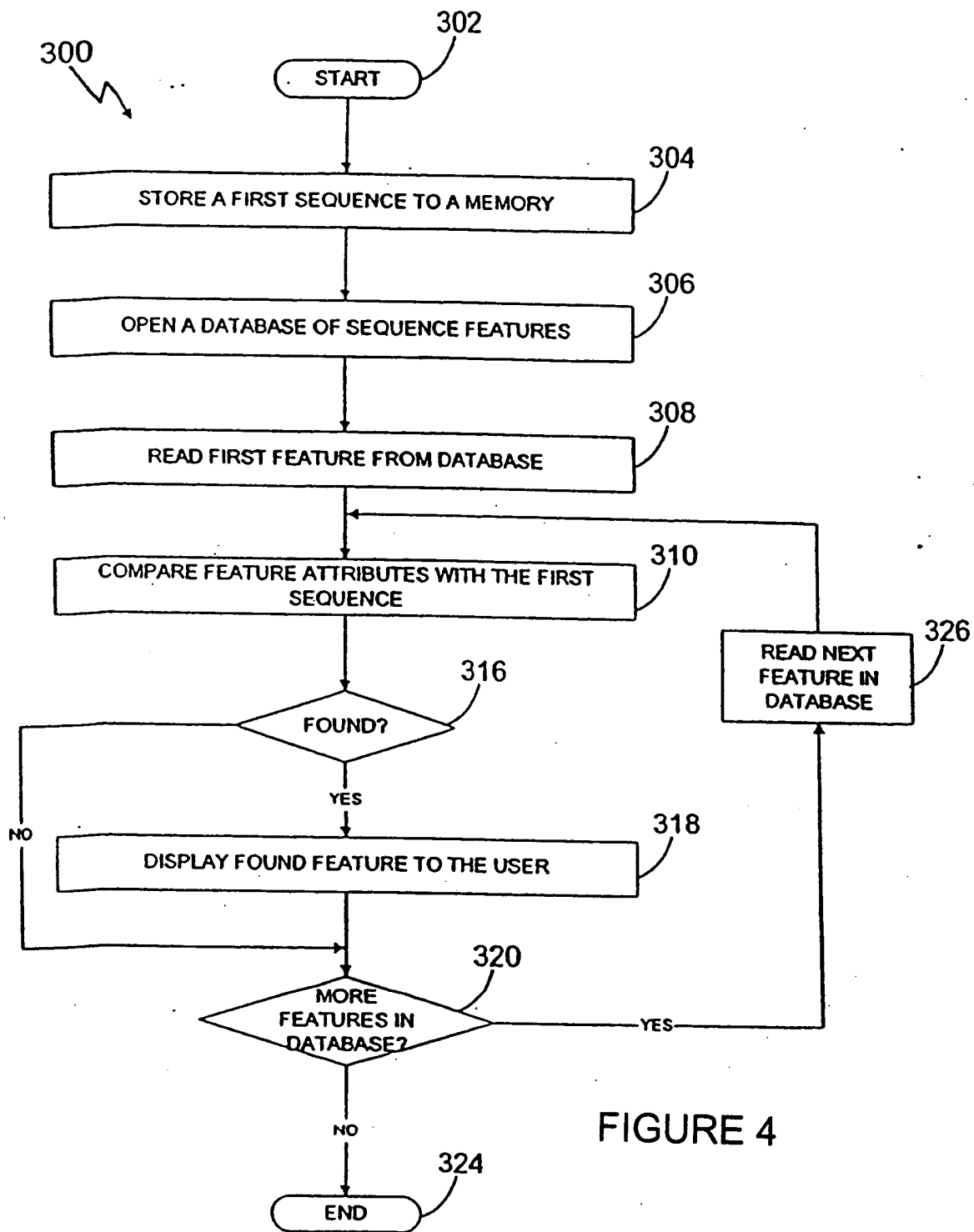


FIGURE 4

# Properties of Diversa Fluorescent Proteins

## DVSACyan

Number of amino acids	227	253
Calculated subunit mass (kDa)	25.9	28.6
Total mass (kDa)	51.8	57.3
Excitation maximum (nm)	448(463)	487
Emission maximum (nm)	491	507
Quantum yield	0.76	0.61
Extinction coefficient ( $M^{-1} \text{ cm}^{-1}$ )	18,900	98,200

Figure 5

DIVERSA

# DVSAGreen vs. Other GFPs

## Excitation Maxima

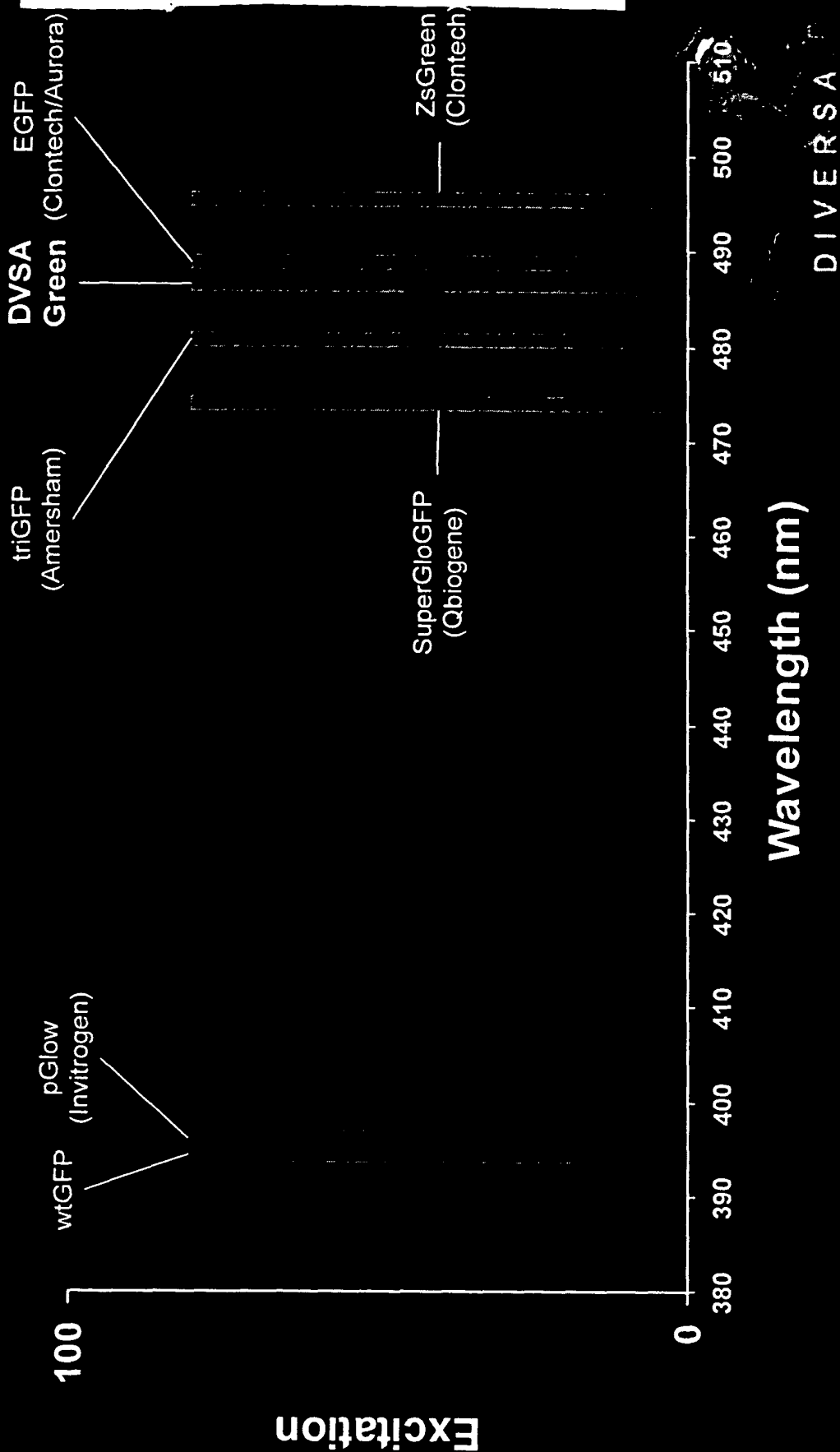


Figure 6

# DVSAGreen vs. Other GFPs

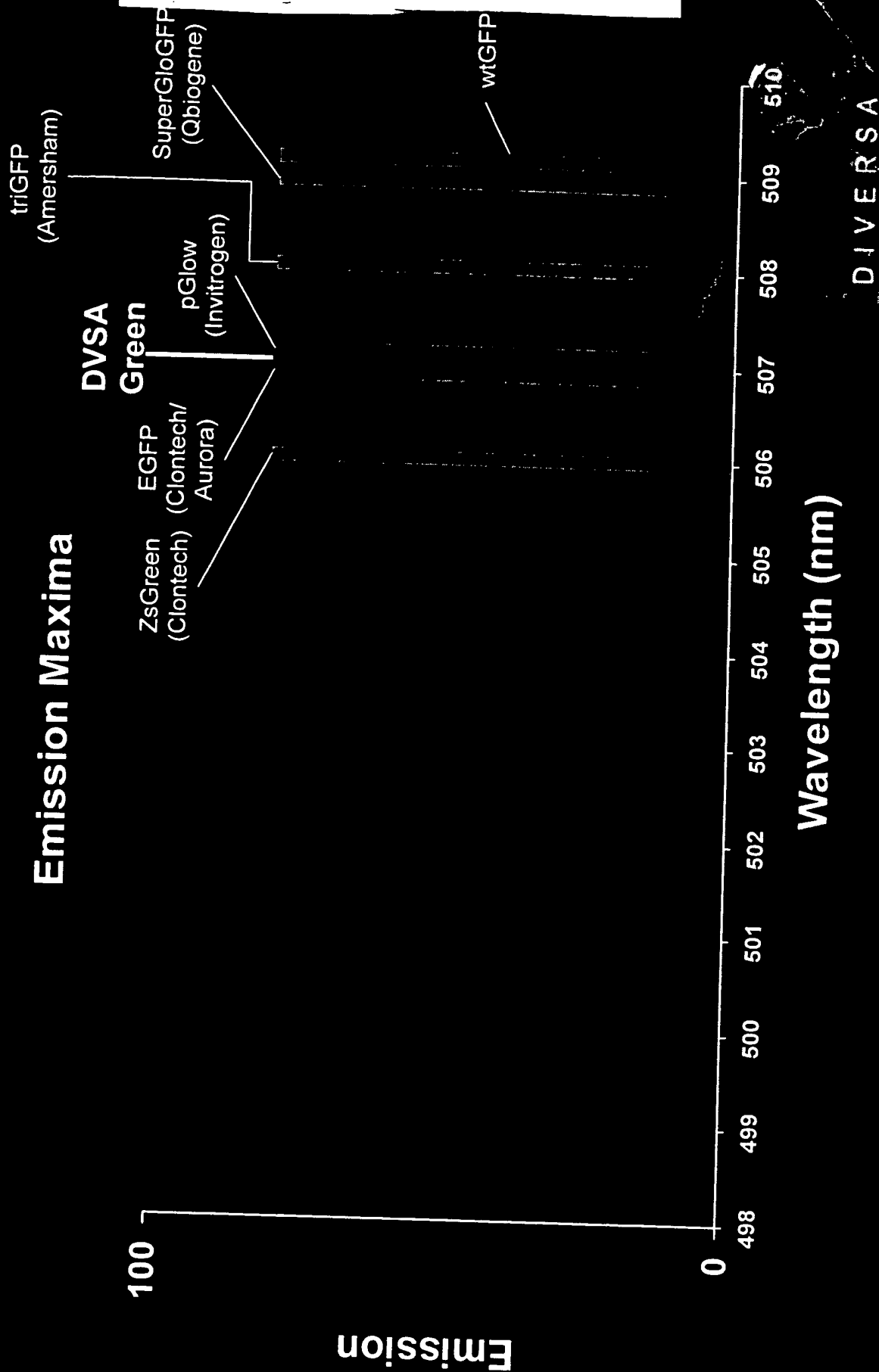


Figure 7

# DVSACyan vs. Other Blue/Cyan FPs

## Excitation Maxima

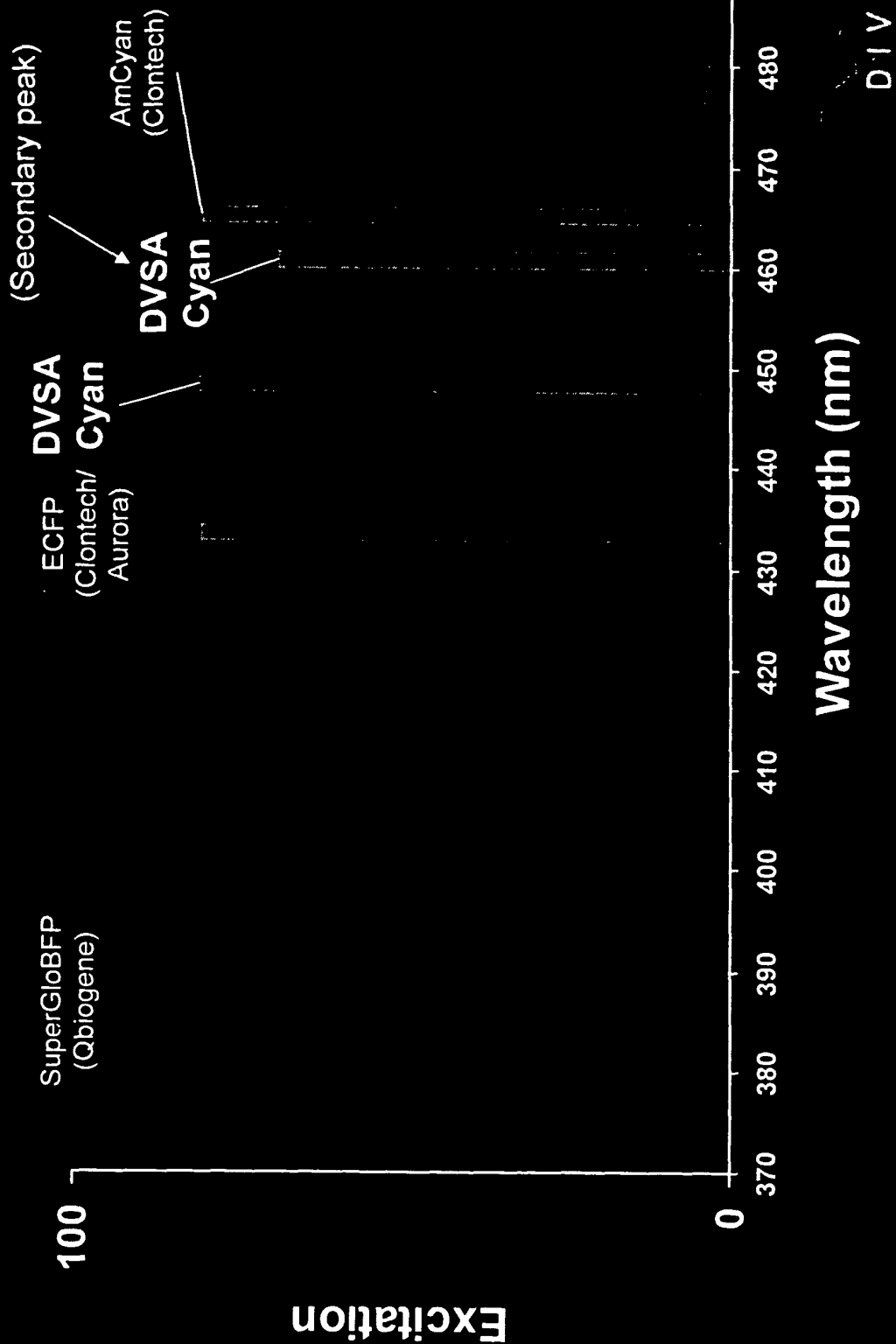


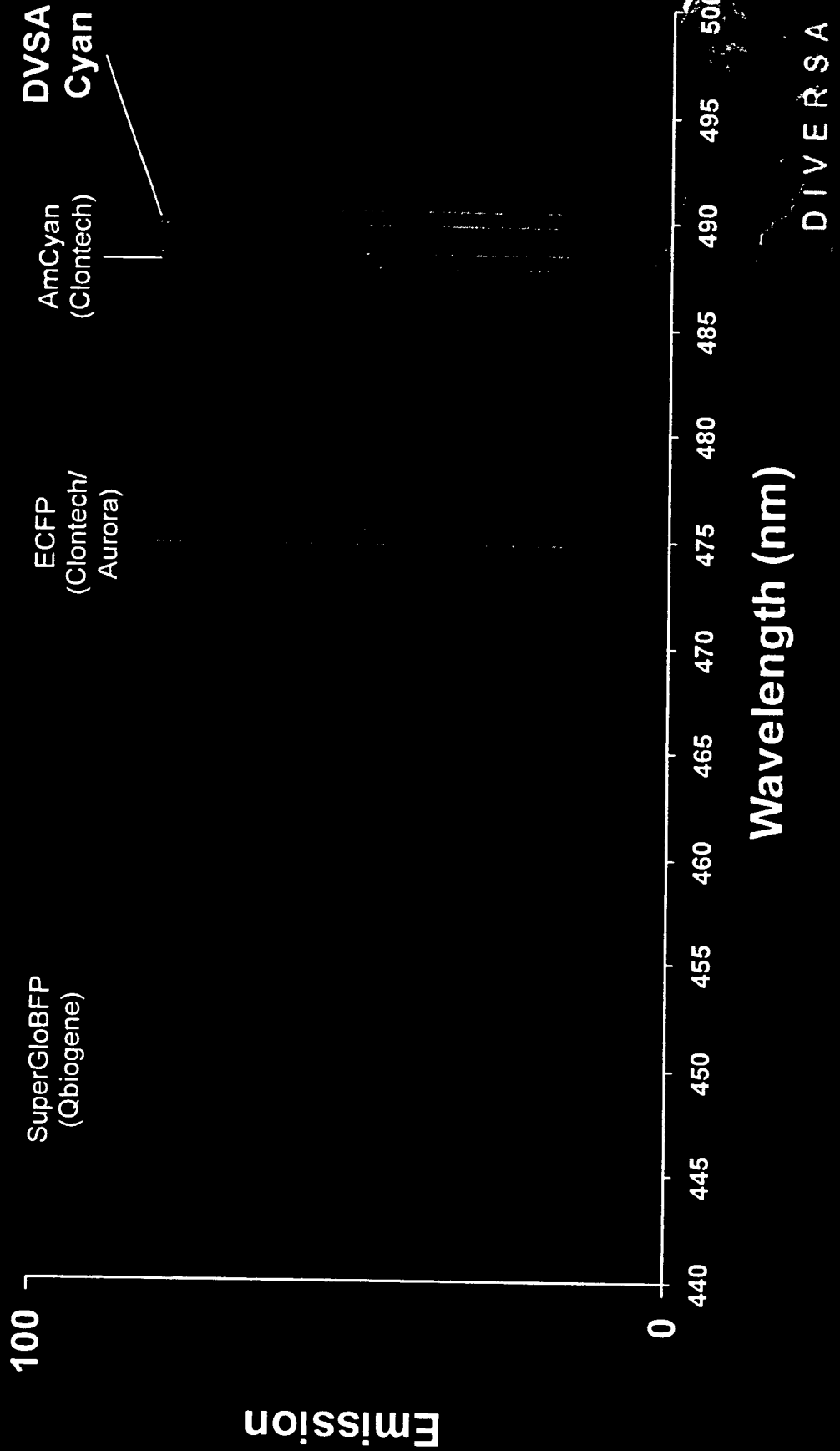
Figure 8



Figure 9

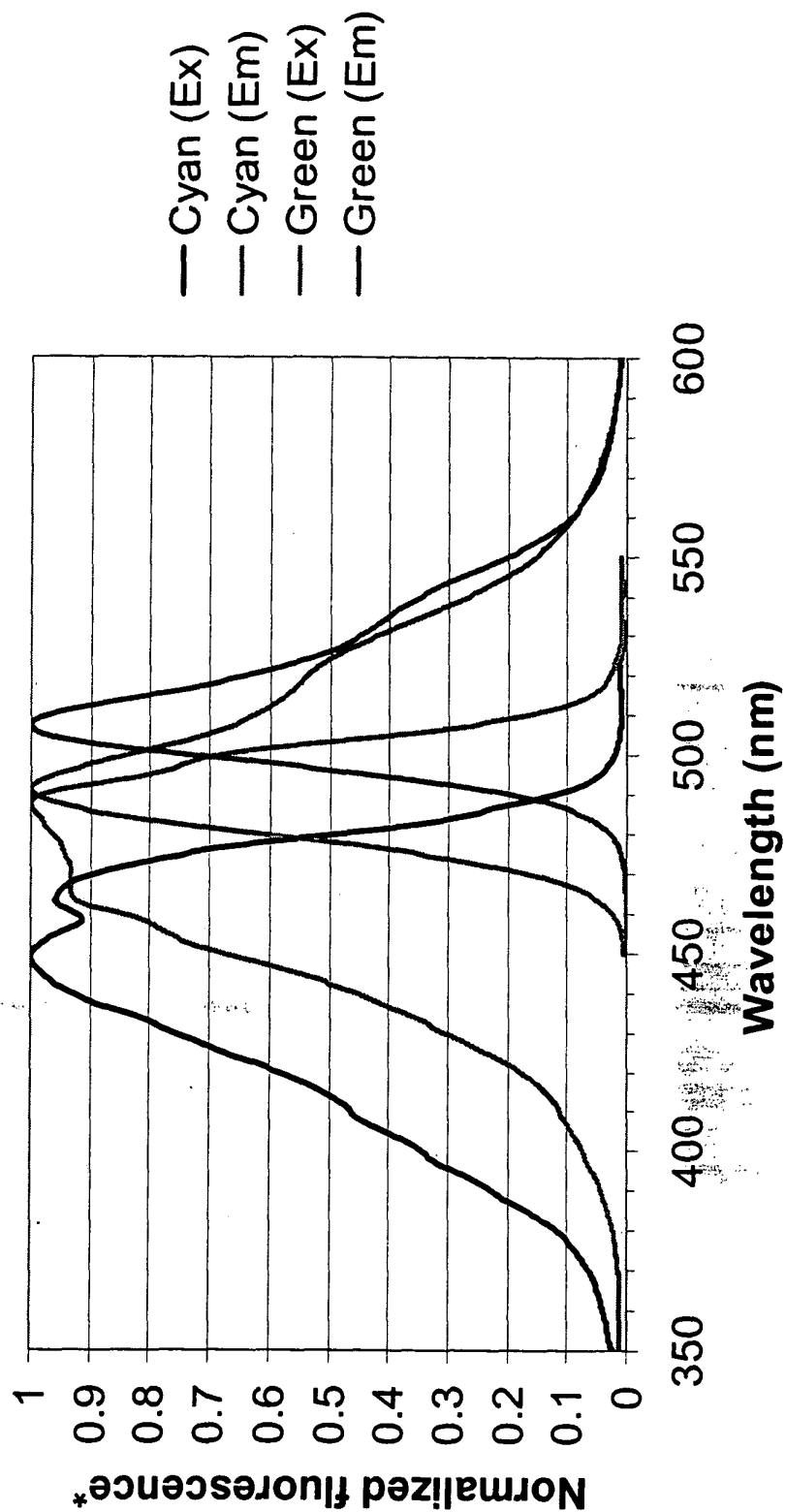
# DVSA Cyan vs. Other Blue/Cyan FPs

## Emission Maxima



# Excitation and Emission Spectra

## Diversa Fluorescent Proteins



\*Spectra normalized to the peak excitation and emission fluorescence for each protein

DIVERSA

Figure 10

# DVSA Green protein is brighter than EGFP

Quantum yield      Extinction coefficient      Relative brightness\*

( $M^{-1} \text{ cm}^{-1}$ )

wtGFP	0.77-0.80 <sup>1,2</sup>	21,600-27,600 <sup>1,2</sup>	1
EGFP	0.6-0.7 <sup>3,4</sup>	39,200-55,900 <sup>3,4</sup>	1.42-1.77
DVSA Green	0.79	90,000	2.7-3.6
AmCyan	0.24 <sup>2</sup>	40,000 <sup>2</sup>	0.43-0.58
DVSACyan	0.76	18,900	0.65-0.88

\* Relative brightness (maximal extinction coefficient multiplied by quantum yield) as compared to wtGFP

<sup>1</sup> Taken from Heim and Tsien, Current Biology 1996

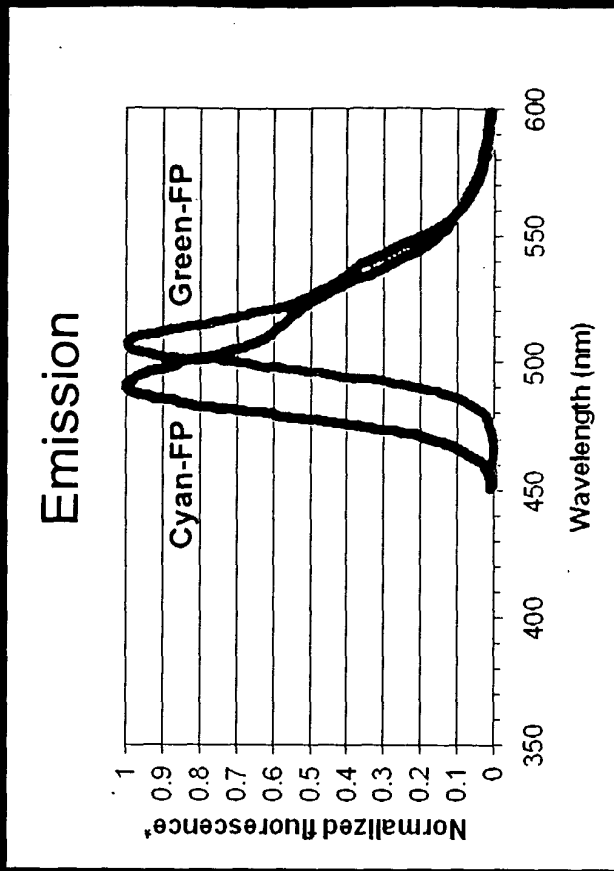
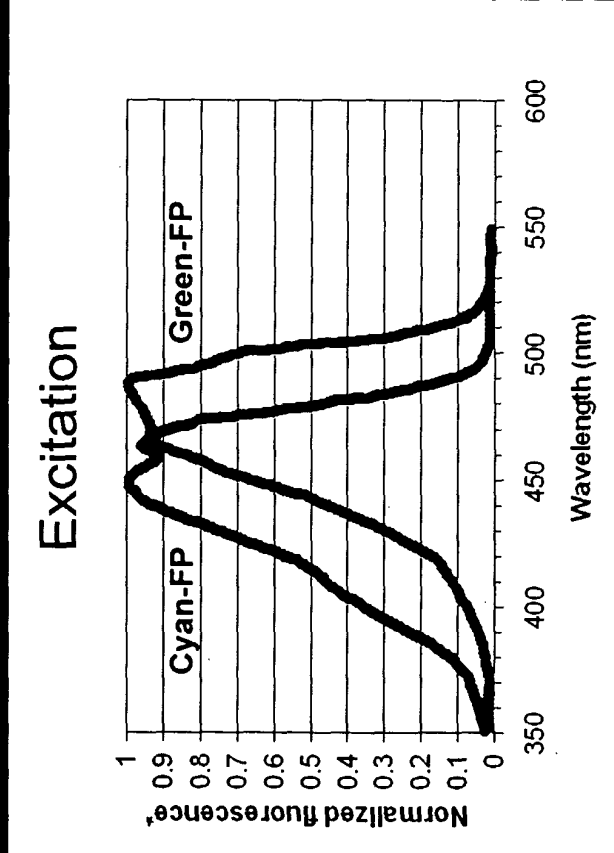
<sup>2</sup> Taken from Matz et al, Nature Biotechnology, 1999

<sup>3</sup> Taken from Zimmer, Chemical Reviews, 2002

<sup>4</sup> Taken from Remington, Nature Biotechnology, 2002

D I V E R S A

# Excitation and Emission Spectra



\*Spectra normalized to the peak excitation and emission fluorescence for each protein

Figure 12

# Relative Brightness

Comparison with Commercially Available Fluorescent Proteins			
	Quantum Yield	Extinction Coefficient (M <sup>-1</sup> cm <sup>-1</sup> )	Relative Brightness*
Discovery Point <sup>1</sup> (wtAvGFP)	0.61	98,200 <sup>2</sup>	2.7-3.6
Wild type AvGFP	0.77-0.80 <sup>1,2</sup>	21,600-27,600 <sup>1,2</sup>	1
EGFP	0.6-0.7 <sup>3,4</sup>	39,200-55,900 <sup>3,4</sup>	1.42-1.77
pGlow	0.79 <sup>3</sup>	30,000 <sup>3</sup>	1.1-1.4
Discovery Point <sup>1</sup> (AmCyan)	0.76	18,900 <sup>2</sup>	0.65-0.88
AmCyan	0.24 <sup>2</sup>	40,000 <sup>2</sup>	0.43-0.58
ECFP	0.4 <sup>3</sup>	32,500 <sup>3</sup>	0.59-0.78
* Relative brightness (maximal extinction coefficient multiplied by quantum yield) as compared to wtAvGFP, + Measured per chromophore			

1. Heim and Tsien. Current Biology 1996
2. Matz et al. Nature Biotechnology. 1999
3. Zimmer. Chemical Reviews. 2002
4. Remington. Nature Biotechnology. 2002

Figure 13

# Summary

## Summary of Diversa's DiscoveryPoint™ Fluorescent Proteins

	DiscoveryPoint™ Green-PP	DiscoveryPoint™ Cyan-PP
Excitation/Emission max (nm)	487/507	448(463)/491
Stoke's shift (nm)	20	43(28)
Maturation time	Within 1 hour	Within 1 hour
Quantum yield	0.61	0.76
Extinction coefficient (M <sup>-1</sup> cm <sup>-1</sup> )	98,200	18,900
Thermostable to 80°C	Yes	Yes
# of amino acids	228	227
Calculated subunit mass (kDa)	26.0	25.9
Total mass (kDa) - dimers	52.0	51.8

Figure 14

DIVERSA

SEQ ID NO: 27 Nucleotide location of segment (start- stop)	SEQ ID NO: 29 Nucleotide location of segment (start- stop)	SEQ ID NO: 31 Nucleotide location of segment (start- stop)
1-53	1-41	1-43
Overhangs on start-stop start-GGA/CCT	Overhangs on start-stop start-GGA/CCT	Overhangs on start-stop start-CATA/GTAT
		CATA/GTAT-TCCT/AGGA
57-78	45-66	97-142
82-116	70-104	TCCT/AGGA-GGA/CCT
120-157	108-145	GGA/CCT-TTT/AAA
161-185	149-173	TTT/AAA-AGG/TCC
190-224	178-212	AGG/TCC-CTC/GAG
228-275	216-266	CTC/GAG-ACCA/TGGT
278-323	269-314	ACCA/TGGT-CCC/GGG
327-354	318-342	CCC/GGG-CT/GA
358-393	346-378	CT/GA-AAG/TTC
397-436	382-421	AAG/TTC-TTC/AAG
441-477	426-471	TTC/AAG-CCT/GGA
481-500	475-504	CCT/GGA-CATC/GTAG
503-542	507-546	CATC/GTAG-GGA/CCT
546-593	550-597	GGA/CCT-GG/CC
596-638	600-639	GG/CC-AAG/TTC
642-end	643-end	AAG/TTC-GA/CT
		GA/CT-GAG/CTC
		GAG/CTC-end

Figure 15